PATENT COOPERATION TREATY

PCT

INTERNATIONAL PRELIMINARY REPORT ON PATENTABILITY

(Chapter II of the Patent Cooperation Treaty)

(PCT Artcle 36 and Rule 70)

					
Applicant's or agent's file reference PCT-2758	FOR FURTHER ACTIO	N	See Form PCT/IPE.	A/416	
International application No.	International filing date(day/r	nonth/year)	Priority date (day/mor	nth/year)	
PCT/KR2005/000235 27 JANUARY 2		7.01.2005)	30 JANUARY 2004 (30.01.2004)		
International Patent Classification (IPC) or national classification and I	PC	•		
C12N 9/24(2006.01)i, C12N 9/9/00(2006.01)i	/30(2006.01)i, C12N 15/50	6(2006.01)i, C1	2N 15/63(2006.01)	i, C12N	
Applicant					
LIFENZA CO., LTD. et al				· · · · · · · · · · · · · · · · · · ·	
1. This report is the international production Authority under Article 35 and to		-	nternational Preliminary	Examining	
2. This REPORT consists of a total	of 4 sheets, incl	uding this cover sh	eet.		
3. This report is also accompanied a. (sent to the applicant an	by ANNEXES, comprising: d to the International Bureau) a	total of7	sheets, as follows	S:	
sheets of the des	scription, claims and/or drawing ntaining rectifications authorized nstructions).	•		_	
beyond the discle Supplemental Bo b. (sent to the International	al Bureau only) a total of (indica	ation as filed, as inc ate type and number	licated in item 4 of Box of electronic carrier(s)	No. I and the	
0 2	isting and/or tables related there be Listing (see Section 802 of the	-			
4. This report contains indications r		•			
Box No. I Basis of the	e report				
	Box No. II Priority				
Box No. III Non-establishment of opinion with regard to novelty, inventive step and industrial applicability					
	Box No. IV Lack of unity of invention				
Box No. V Reasoned citations an	Box No. V Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement				
Box No. VI Certain documents cited				•	
Box No. VII Certain def	Box No. VII Certain defects in the international application				
Box No. VIII Certain observations on the international application					
Date of submission of the demamd	Dat	e of completion of	this report		
		•	•		
24 AUGUST 2005 (24.08.2005)	16 MAY 200	6 (16.05.2006)		
Name and mailing address of the IPEA	/KR Aut	thorized officer	·		
Korean Intellectual Property 920 Dunsan-dong, Seo-gu, Republic of Korea		CHO, YOUNG GYUN			
Facsimile No. 82-42-472-7140	Tel	Telephone No. 82-42-481-8132			

INTERNATIONAL PRELIMINARY REPORT ON PATENTABILITY

International application No.

PCT/KR2005/000235

Box	No.	I Basis of the report	· · · · · · · · · · · · · · · · · · ·	
1.		th regard to the language, this report is based on the erwise indicated under this item. This report is based on translations from the original which is the language of a translation furnished international search (under Rules 12.3 and publication of the international application international preliminary examination (under Rules 12.3).	ginal language into the following lan for the purposes of: d 23.1(b)) n (under Rule 12.4)	
	to the	regard to the elements of the international applicate receiving Office in response to an invitation under the second seco	ranished received by this Authority on received by this Authority on	·
	× · · · · · · · · · · · · · · · · · · ·	pages*	received by this Authority on	as originally filed/furnished
3.	· .	The amendments have resulted in the cancellation the description, pages the claims, Nos. the drawings, sheets the sequence listing (specify): any table(s) related to sequence listing (specify)		
4. [This report has been established as if (some of) the made, since they have been considered to go beyo (Rule 70.2(c)). the description, pages the claims, Nos. the drawings, sheets the sequence listing (specify): any table(s) related to sequence listing (specify)	end the disclosure as filed, as indicated	d in the Supplemental Box
* If	item	4 applies, some or all of those sheets may be mark	ed "superseded."	

INTERNATIONAL PRELIMINARY REPORT ON PATENTABILITY

International application No.

PCT/KR2005/000235

Box No. V Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement

Statement			
Novelty (N)	Claims	1-10	YES
	Claims	None	NO
Inventive step (IS)	Claims	1-10	YES
	Claims	None	NO
Industrial applicability (IA)	Claims	1-10	YE
	Claims	None	, NO

2. Citations and explanations (Rule 70.7)

The following documents have been considered for the purpose of this report:

D1: WO 2003/018790 A1 (LIFENZA CO., LTD.) 6 MARCH 2003

D2: WO 2001/066570 A1 (KIM et al.) 13 SEPTEMBER 2001

D3: J. Microbiol. Biotechnol., Vol. 9(3), pp. 260-264 (1999)

D4: Biosci. Biotechnol. Biochem., Vol. 64(2), pp. 223-228 (2000)

The present invention relates to an enzyme, having the amino acid sequence of SEQ. ID. NO:1, with the activity of hydrolyzing amylopectin, starch, glycogen and amylose; a gene (SEQ. ID. NO:2) encoding said enzyme; a transformed cell expressing said gene; a method of producing said enzyme; and a composition for the dextran removal and the plaque elimination.

D1-D4 disclose the DEXAMmase (dextranase and amylase), having antiplaque and anticaries activities, having dextranase and amylase activities simultaneously and degrading insoluble glucans, from *Lipomyces starkeyi* KSM 22 (KFCC 11077); a preparation method of DEXAMase; and an oral composition comprising the same.

However, none of the prior art documents disclose the amino acid sequence of the enzyme (SEQ. ID. NO:1) capable of hydrolyzing amylopectin, starch, glycogen and amylose, and the nucleotide sequence of gene (SEQ. ID. NO:2) encoding the enzyme, and said enzyme in this invention cannot be derived in an obvious manner from the prior art documents.

Therefore, claims 1-10 meet the requirements of novelty, inventive step and industrial applicability under PCT Article 33(2)-(4). //

INTERNATIONAL PRELIMINARY REPORT ON PATENTABILITY

International application No.

PCT/KR2005/000235

Supplemental Box Relating to Sequence Listing						
Continuation of Box No. I, item 2:						
1. With regard to any nucleotide and/or amino acid sequence disclosed in the international application and necessary to the claimed invention, this report was established on the basis of:						
a. type of material a sequence listing table(s) related to the sequence listing						
b. format of material on paper in electronic form						
c. time of filing/furnishing contained in the international application as filed filed together with the international application in electronic form furnished subsequently to this Authority for the purposes of search and/or examination received by this Authority as an amendment* on 07/04/2006						
2. In addition, in the case that more than one version or copy of a sequence listing and/or table(s) relating thereto has been filed of furnished, the required statements that the information in the subsequent or additional copies is identical to that in the application as filed or does not go beyond the application as filed, as appropriate, were furnished.						
3. Additional comments:						

10/588052 PEA/KR 07.04.2006. AP20 Rec'd PCT/PTO 31 JUL 2006

maltodextrin (Mn), before and after being hydrolyzed by the (lanes 1 and 2 in panel A, respectively) enzyme and maltooligosaccharide samples (1% w/v) are analyzed after allowed to purified LSA is react with a series of maltooligosaccharides including G1 (glucose) G7 to (maltoheptaose) (lanes 1 to 7 in panel B, respectively).

DESCRIPTION OF THE PREFERRED EMBODIMENTS

10

15

20

25

The acquisition of a gene coding for the carbohydrolase (LSA) of the present invention starts by culturing Lipomyces starkeyi in a medium containing starch. Next, on the basis of N-terminal amino acid sequences of carbohydrate hydrolyzing enzymes purified from L. starkeyi, primers comprising expected conserved regions are constructed, followed by PCR with the primers. The PCR product, approximately 2 kb long, is used for 5' RACE and 3' RACE to allow for a complete carbohydrolase gene (LSA). After being amplified by PCR, the gene is cloned in the vector pRSETB (Invitrogen, U.S.A.) with which Escherichia coli DH5@/pRLSA is then transformed.

 $\it L.~starkeyi$ is known to produce endo-dextranase (EC 3.2.1.11) which degrades dextran and α -amylase which degrades starch. This microorganism has been applied to foods and not yet reported to produce antibiotics or other toxic metabolites.

Most of the dextranases produced by microorganisms, except for a few derived from bacteria, are known as inducible enzymes. L. starkeyi ATCC74054, reported first in U.S. Pat.

AMENDED SHEET (ART. 34)

No. 5,229,277, produces both dextranase and amylase whose characteristics are also disclosed. It is also reported that the strain produces low molecular weight dextrans from sucrose and starch. On the basis of the findings, the present inventors have acquired Korean Pat. No. 10-0358376 on Oct. 11, 2002 (corresponding to U.S. Pat. No. 6,485,953 dated Nov. 26, 2002) which relates to a DXAMase enzyme capable of hydrolyzing both dextran and starch, a microorganism producing the enzyme (identified as *Lipomyces starkeyi* KFCC-11077), and a composition comprising the enzyme.

10

15

20

25

The enzyme expressed from the gene (*lsa*) of the present invention is a carbohydrolase capable of hydrolyzing amylopectin, starch, glycogen and amylose. Also, the enzyme according to the present invention is found to degrade dextran, alpha-cyclodextrin and pullulan. The enzyme is highly stable. Not only is its activity 90% of its maximum over a relatively broad pH range (pH 5-8), but also it is not inhibited even by a denaturation solution such as an EGTA-containing solution. Ca²⁺ or Mg²⁺ serves as a cofactor for the enzyme.

Also, the present invention is directed to a novel microorganism carrying the gene coding for the carbohydrolase. The strain *Escherichia coli* DH5@/pRLSA according to the present invention was deposited in the Korean Collection for Type Cultures (KCTC) located in Yusung Gu, Daejeon City, South Korea, with the accession number of KCTC 10573BP, on Dec. 24, 2003.

Also, the present invention is directed to a method of



producing the carbohydrolase. First, the strain *Escherichia* coli DH50/pRLSA is cultured. After being harvested from the culture, the cells are disrupted using glass beads to isolate the carbohydrolase therefrom.

A composition comprising the enzyme of the present invention may be used in a variety of oral care applications. By virtue of its ability to degrade polysaccharides such as dextran and amylose, the enzyme of the present invention is also effectively used to remove dextran during sugar production. Additionally, compositions comprising the enzyme according to the present invention can be applied to foods such as gum, drinks, milks, etc. and their constituents may be readily determined by those who are skilled in the art.

A better understanding of the present invention may be obtained through the following examples which are set forth to illustrate, but are not to be construed as the limit of the present invention.

EXAMPLE 1: 1sa gene cloning in Lipomyces starkeyi

20

25

5

10

15

1) Strain and plasmid

Lipomyces starkeyi KFCC 11077, which produces DXAMase having dextranase and amylase activity, was used as a DNA donor for cDNA isolation and amylase gene selection. General DNA manipulation and DNA sequencing were carried out with Escherichia coli DH5 α and pGEM-T easy (Promega, USA). For the construction of a cDNA library, E. coli XL1-Blue and SOLR (Stratagene, USA) were used as host cells with lambda phase



primer 5'-CTCTACATGGAGCAGATTCCA-3' which respectively correspond to N-terminal and C-terminal amino acid sequences of the protein showing dextranase and amylase characteristics. After being separated on agarose gel, the PCR product was purified with an AccPrepTM gel extraction kit (Bioneer, Korea) and ligated with pGEM-T easy vector (Promega, USA). Base sequencing was performed using ABI PRISM Cycle Sequencing Kit (Perkin Elmer Corp. USA) in a GeneAmP 9600 thermal cycler DNA sequencing system (Model 373-18, Applied Biosystems, USA).

10

15

20

25

8) Heterologous expression and purification of LSA protein in *E. coli*

The gene *lsa* was inserted into the SacI-EcoRI site of pRSETB vector (Invitrogen USA) to prepare a recombinant vector pRSET-LSA. Escherichia coli DH50/pRLSA transformed with pRSET-LSA was cultured at 37°C to a midstationary phase in an LB medium containing 50 mg/l ampicillin. After the addition of the culture to a final concentration of 1 mM, IPTG incubation was carried out at 28°C for 6 hours. Cells were harvested by centrifugation (5000 g \times 10 min), washed with 0.1 M potassium phosphate (pH 7.4 and lyzed by sonication. Purification of the expressed protein was performed with Ni^{2+} nitrilotriacetic acid-agarose (NTA) (Quiagene, Germany). cell lysate was combined with Ni²⁺-NTA and allowed to stand for 1 hour at 4°C, and the mixture was loaded onto a column which was then washed four times with a washing buffer. Each 0.5 ml of the protein fraction was emulsified with a buffer.

WHAT IS CLAIMED IS:

- 1. A protein, comprising an amino acid sequence of SEQ. ID. No. 1, which has the activity of hydrolyzing amylopectin, starch, glycogen and amylose, a derivative thereof, or a fragment thereof.
- 2. A gene of SEQ. ID. No. 2, encoding the protein, the derivative, or the fragment of claim 1, a derivative thereof, or a fragment thereof.
 - 3. A transformed cell, expressing the gene, the derivative, or the fragment of claim 2.
- 4. The transformed cell as defined in claim 2, wherein the cell is prokaryotic or eukaryotic.
- 5. The transformed cell as defined in claim 3 or 4, wherein the cell is *Escherichia coli* DH5@/pRLSA deposited with the accession number of KCTC 10573BP.
 - .6. A method of producing an enzyme having activity of hydrolyzing amylopectin, starch, glycogen and amylose, comprising:
- culturing the cell of claim 3;
 expressing the enzyme in the cultured cell; and
 purifying the expressed enzyme.



[Sequence Listing]

· <110> Lifenza Co., Ltd. ` 5 <120> PROTEIN WITH ACTIVITY OF HYDROLYZING AMYLOPECTIN, STARCH, GLYCOGEN AND AMYLOSE, GENE ENCODING THE SAME, CELL EXPRESSING THE SAME, AND PRODUCTION METHOD THEREOF <150> KR2004-0006186 <151> 2004-01-30 10 <160> <170> KopatentIn 1.71 15 <210> <211> 647 <212> PRT <213> Artificial Sequence 20 <220> Escherichia coli DH5@/pRLSA <400> · 1 25 Met Leu Leu IIe Asn Phe Phe IIe Ala Val Leu Gly Val IIe Ser Leu 15 . 10 Ser Pro IIe Val Val Ala Arg Tyr IIe Leu Arg Arg Asp Cys Thr Thr 30 20 30 Val Thr Val Leu Ser Ser Pro Glu Ser Val Thr Ser Ser Asn His Val 35 40 45 35 Glu Leu Ala Ser His Glu Met Cys Asp Ser Thr Leu Ser Ala Ser Leu 50 55 60

AMENDED SHEET (ART. 34)

	011	10.40					•
	211>	1946					
	<212>	ONA .	•				
5	213 >	Artificial Sec	quence				
	<220>						
	<223> E	scherichi	a coli	DH5@/pH	RLSA		
10							
	<400>	2				•	
*		a tcaactttt c	ratroctott	ctoopantoa	tateactete	toctattoto	60
	argrigorg.		arogorgrr	Cryyyayrya	tattactgto	toctatigig	00
	attactact	· atattattaa a	vocačat tao	antanatta	annint tata	0100001000	100
	grigorogi	atattcttcg a	icgaga i igc	actacagita	cggictigic	ctccctgag	120
15				•			
	tetgtgaega	a gttcgaacca t	gttcagcta	gccagtcatg	agatgtgcga	cagtaccttg	180
		•			· .		
	tcagcgtcc	tttatatcta c	aatgatgat	tatgataaga	ttgtgacact	ttattatctt	240
20	acatcgtcgg	gcacaactgg g	tccgtaaca	gcgtcttatt	cttctagttt	gagtaacaac	300
					•		
	tgggaattgi	ggtctctctc g	gctccggct	gcagatgctg	tcgagatcac	tggagctagt	360
	tatgtagaca	gcgatgcatc t	gcgacatac	accacatett	ttoatatacc	tcttactacc	420
25							120
_•	acnacaacot	cgtcgtcttc t	actaataca	acttcaacat	ctantctaac	cacaacatot	480
	uoguouuog i	ogrogro, rro r	goragrgog		·	Cacaacatct	400
	aatatttaa	tttooototo	atoootooo				E 40
	agigiticca	tttcggtgtc c	giccciaca	ggaacagctg	caaattggcg	aggtagggct	540
30	atctatcaga	togtgactga ta	agatttgca _.	cgcactgacg	gctccaccac	atatttatgc	600
	gatgttaccg	atagggtcta t	tgcggaggg	tcttatcagg	ggattatcaa	tatgctggat	660
	tacatccaag	gcatgggctt ta	actoctatt	tggatttctc	ctatagtgga	aaatattccc	720
35							
	gatgacaccg	gatacggtta c	gcatatcat	ggttattgga	tgaaagatat	cttcgccctg	780
	_		-	55			

<210>